

SEQUENCE LISTING

<110> Martin Laforest
Nathalie Hubert
Benoit S. Landry

<120> Methods for Relative Quantification of
Specific Nucleic Acid Sequences

<130> 14187-1PCT

<160> 8

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 1600

<212> DNA

<213> E. coli

<400> 1

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<210> 2

<211> 1574

<212> DNA

<213> Brassica napus

<400> 2

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<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Sequence to be used as a primer

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cagctgtgct cgacgttgct a

21

<210> 4

<211> 20

<212> DNA

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<220>

<223> Sequence to be used as a primer

<400> 4

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<210> 5

<211> 21

<212> DNA

<213> Artificial Sequence

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<210> 6
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<212> DNA
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<220>
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<400> 6
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<210> 7
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<400> 7
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<210> 8
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<220>
<223> Sequence to be used as a primer

<400> 8
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<210> 9
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<212> DNA
<213> Artificial Sequence

<220>
<223> This sequence is the dispensation order of
nucleotides given in Example 1 of the application.

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